

SEQ.ST25  
SEQUENCE LISTING

&lt;110&gt; Gonzalgo, Mark L.

Jones, Peter A.

Liang, Gangning

&lt;120&gt; CANCER DIAGNOSTIC METHOD BASED UPON DNA METHYLATION DIFFERENCES

&lt;130&gt; 47675-21

&lt;140&gt; 09/887,941

&lt;141&gt; 2001-06-22

&lt;150&gt; US 09/094,207

&lt;151&gt; 1998-06-09

&lt;160&gt; 17

&lt;170&gt; PatentIn version 3.1

&lt;210&gt; 1

&lt;211&gt; 530

&lt;212&gt; DNA

&lt;213&gt; homo sapiens

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cacggggtag tcggaggggg aggtccatct ccctttccct tgctgtccat ctccacagaa 180  
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gccgcctgtg gcgcccagtc cgaagctgct gttttcagga gggctagtgg gctaagaaaa	420
gagctcaccg actgactgcc caacagctgt tgcgagccag tgctaggctg cagacagcct	480
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gaggtaggga aagcgagctt caccggggcg gctacgatga gtagcatgac gggcagcagc	180
agcagcagcc agcaaaagcc tagcaaagtg tccagctgct gcactgccgc ggggactccc	240
acatcaccat gactagttgt gcaactctgc agcagaaacg gcttccgagg aacacaggat	300
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gaccgaccga gaagaaggga cgtcccgg	148

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&lt;212&gt; DNA

&lt;213&gt; homo sapiens

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&lt;211&gt; 178

&lt;212&gt; DNA

&lt;213&gt; homo sapiens

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<211> 220

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aagattgtag agctagctca cagcggggcc cggccgtgcg acatttcccg aattctgcag 180

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<212> DNA

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<400> 17

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